

# The early marine distribution of Atlantic salmon in the Northeast Atlantic: A genetically informed stock specific synthesis

John Gilbey, Vidar Wennevik, Kjell Utne, Philip McGinnity, Eric Verspoor

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# Introduction

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ORIGINAL ARTICLE

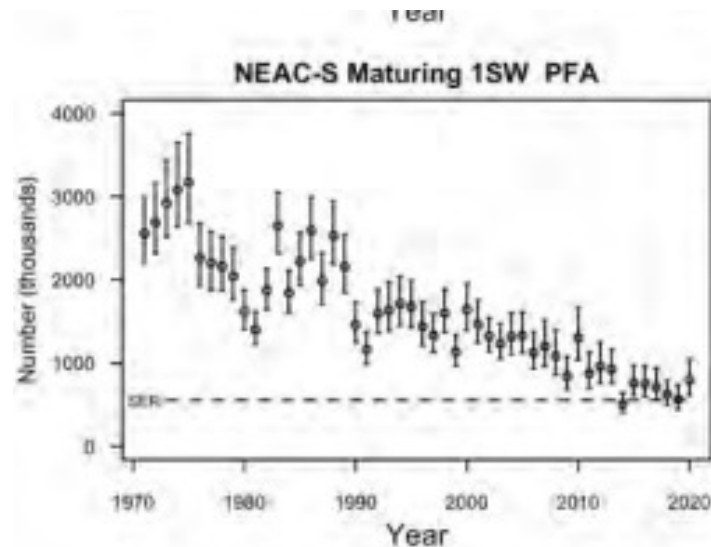
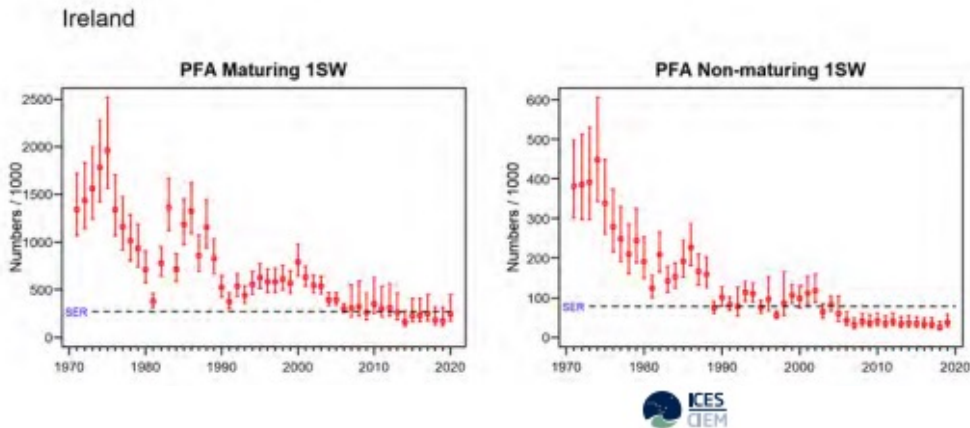


WILEY

## The early marine distribution of Atlantic salmon in the North-east Atlantic: A genetically informed stock-specific synthesis

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# Why increasingly poor survival of Atlantic salmon at sea?



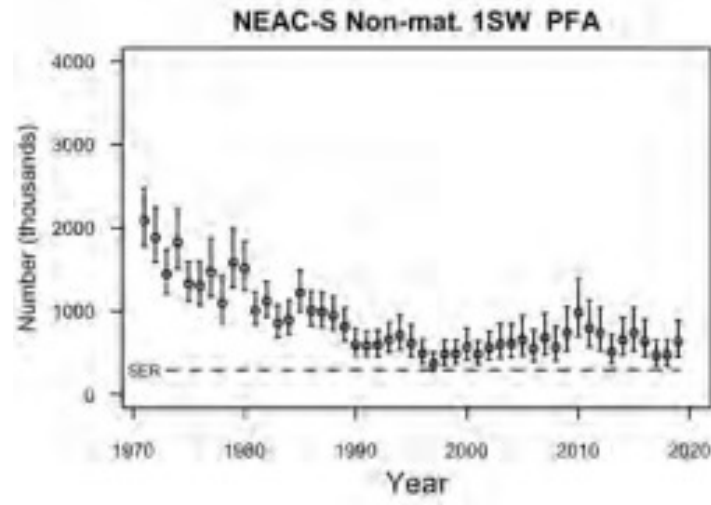
## WORKING GROUP ON NORTH ATLANTIC SALMON (WGNAS)

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RAPPORTS SCIENTIFIQUES DU CEM



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***“Knowing the distribution of marine animals is central to understanding climatic and other environmental influences on population ecology.”***

MacKenzie et al. (2011). Scientific Reports | 1:21 | DOI: [10.1038/srep00021](https://doi.org/10.1038/srep00021)

# Why increased mortality at sea?

To answer question need to take a step back –

Need to describe/define as accurately as possible the theatre (location) of action – or ‘domains’ as Crozier et al. 2018 proposed recently

Provide focus for spatially and temporally targeted analyses of potential explanatory variables - physical, chemical, biological/ecological descriptions of the environment

Move from theoretical ideas about distribution to the real knowledge

Two important initiatives (SALSEA-Merge) more recently (SeaSALAR)

**Concentrate on two areas – early post-smolts and stock specific ID**



## Atlantic salmon mortality at sea: Developing an evidence-based “Likely Suspects” Framework

Walter Crozier, Ken Whelan, Mathieu Buoro, Gerald Chaput, Jason Daniels, Sue Grant, Kim Hyatt, James Irvine, Niall Ó'Maoiléidigh, Etienne Prévost, Etienne Rivot, Ian Russell, Michael Schmidt and Brian Wells



**Why focus on post smolts?**

# Why focus here on post smolts?

Definition of a post smolt? Time of entry into sea to end of calendar year – here **early summer** was the focus - **May, June, July, August**;

Reasonable to presume period of highest vulnerability (mortality) – small animals typically in early summer about 25-35 cm

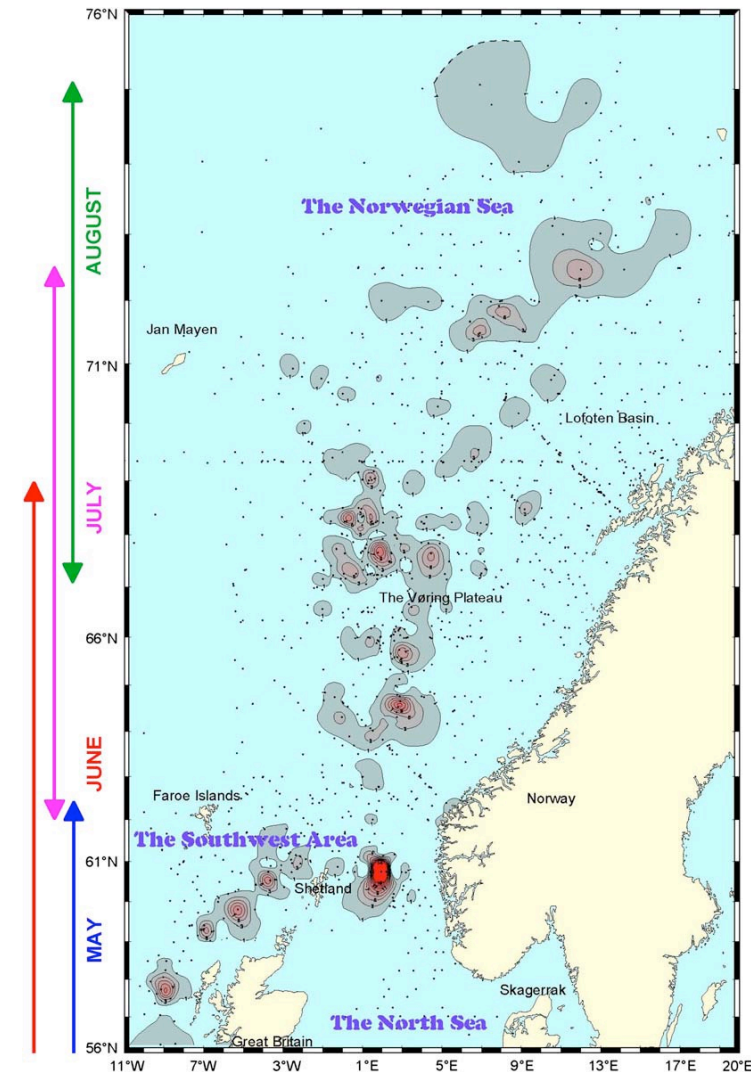
Likely to be heavily influenced by and likely evolved to exploit oceanic conditions - Dingle, H., & Drake, V. A. (2007). What is migration? *BioScience*, 57(2), (e.g. piggy- back ocean currents; Dadswell et al. 2010 – ‘Merry go round hypothesis);

First environments encountered on leaving river (not confounded by multiple environments or life history transitions accumulated by older life history stages) – simpler to get handle on fish ecosystem relationships

Considerable amount of information already known about distribution of summer post smolts in the NE Atlantic

## Feeding of Atlantic salmon (*Salmo salar* L.) post-smolts in the Northeast Atlantic

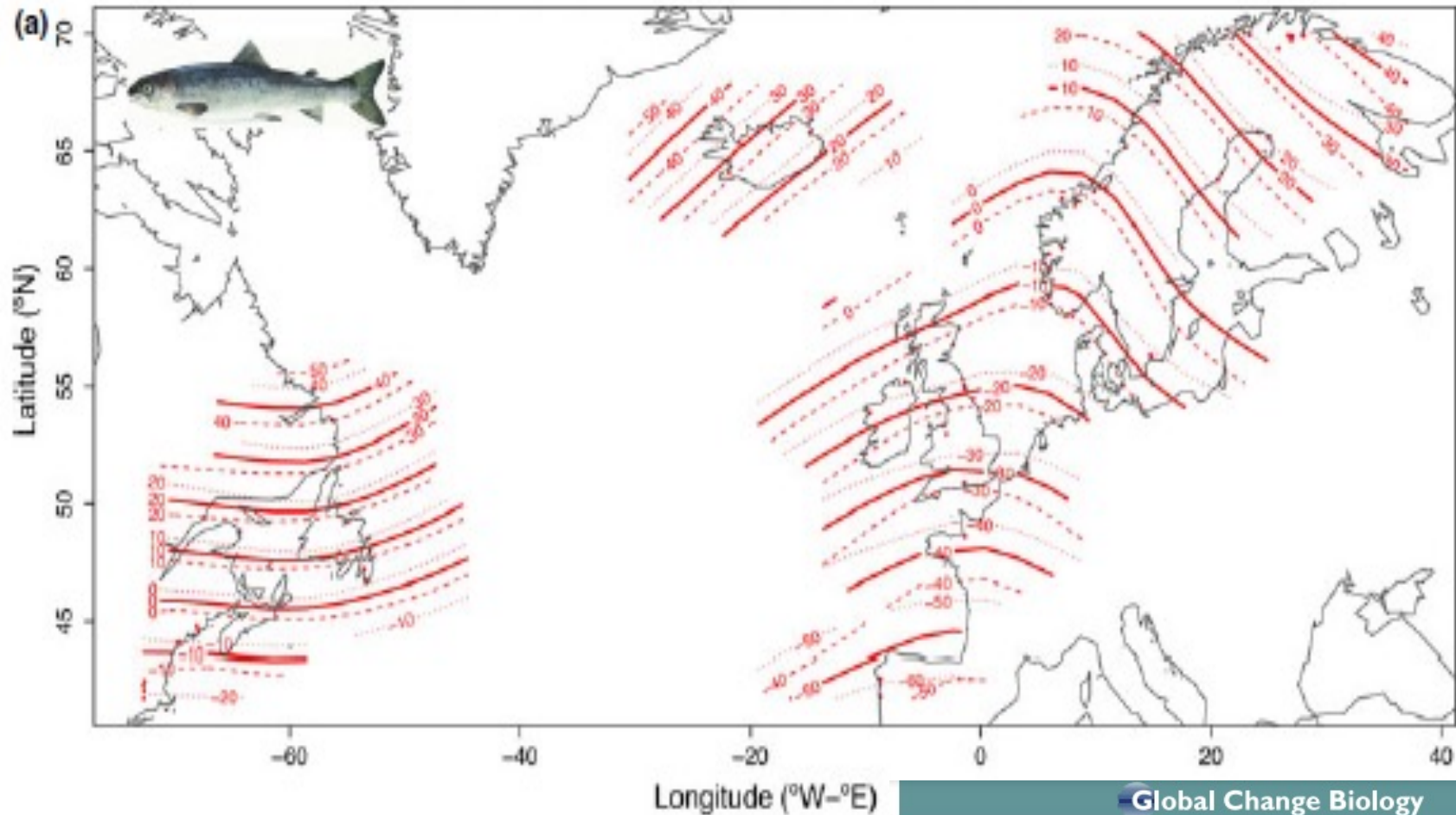
Monika Haugland, Jens Christian Holst, Marianne Holm, and Lars Petter Hansen



**Why stock specific?**



**Geography:** travel distances (35° latitude, ~2,500Km); timing of sea entry (varies by 90 days across NE Atlantic)



Global Change Biology (2014) 20, 61–75, doi: 10.1111/gcb.12363

**Basin-scale phenology and effects of climate variability on global timing of initial seaward migration of Atlantic salmon (*Salmo salar*)**

JAIME OTERO\*, JAN HENNING L'ABÉE-LUND†, TED CASTRO-SANTOS‡,

# **Evolved traits** (important adaptations in contemporary marine environments among individual genetically distinct river populations)

- Phenological – sea entry time
- Morphological (age and size – mouth gape swimming speed and capacity – carry over from river environments)
- Migratory behaviours - sensitivity to magnetic fields, direction finding
- Inherent growth rate
- Metabolic potential (SMR, MMR, AS) – energy acquisition, defence and use
- Immunological
- Predator avoidance - shoaling
- Phylogenetic (legacy – evolved traits to historical environments)



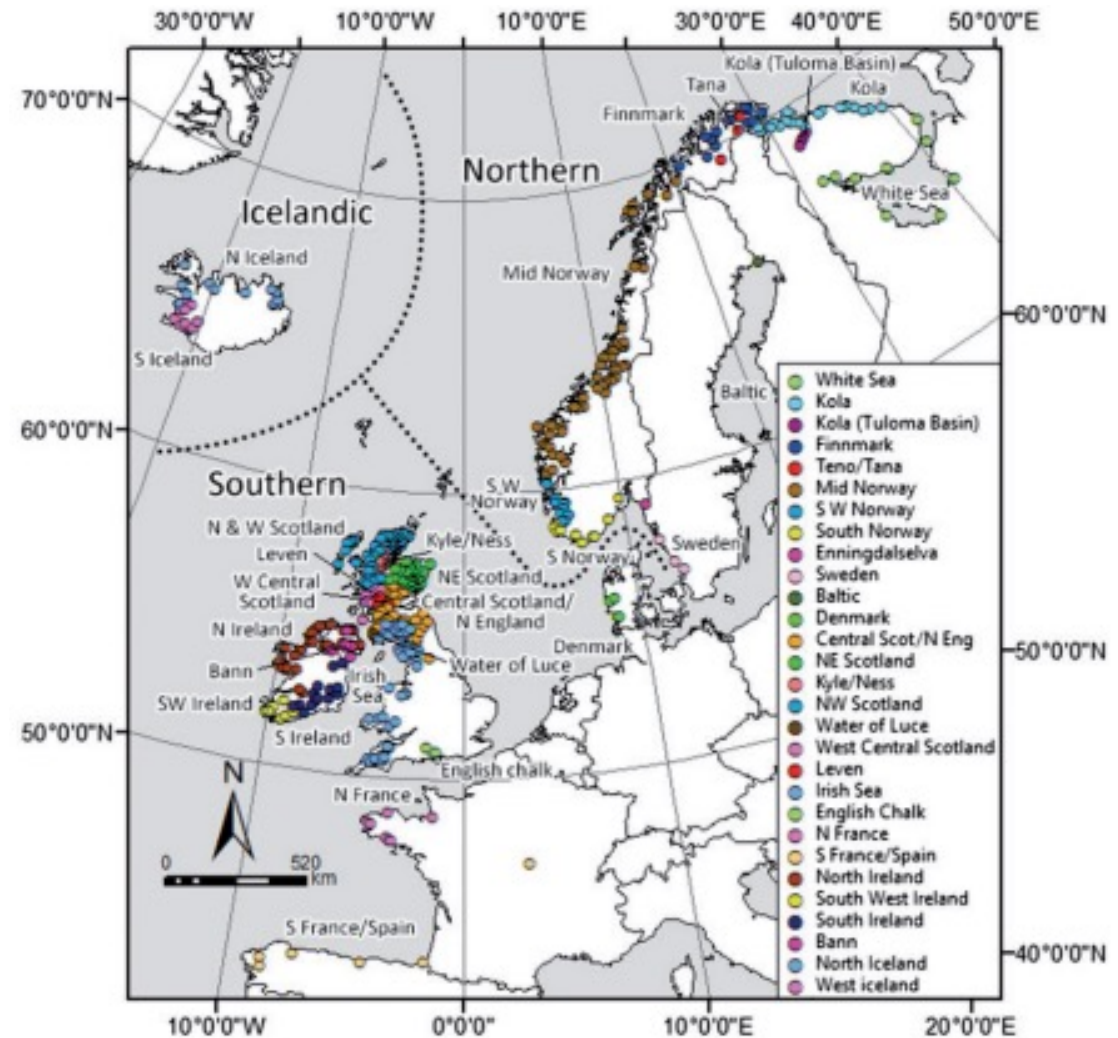
# **Genetic Stock Identification**

# Establishing a baseline (Gilbey et al. 2018)

33,000 individuals

550 sites

325 rivers (Spain to Russia)



ICES Journal of Marine Science



International Council for the Exploration of the Sea  
ICES  
CIEM  
Coastal Information and Monitoring Programme

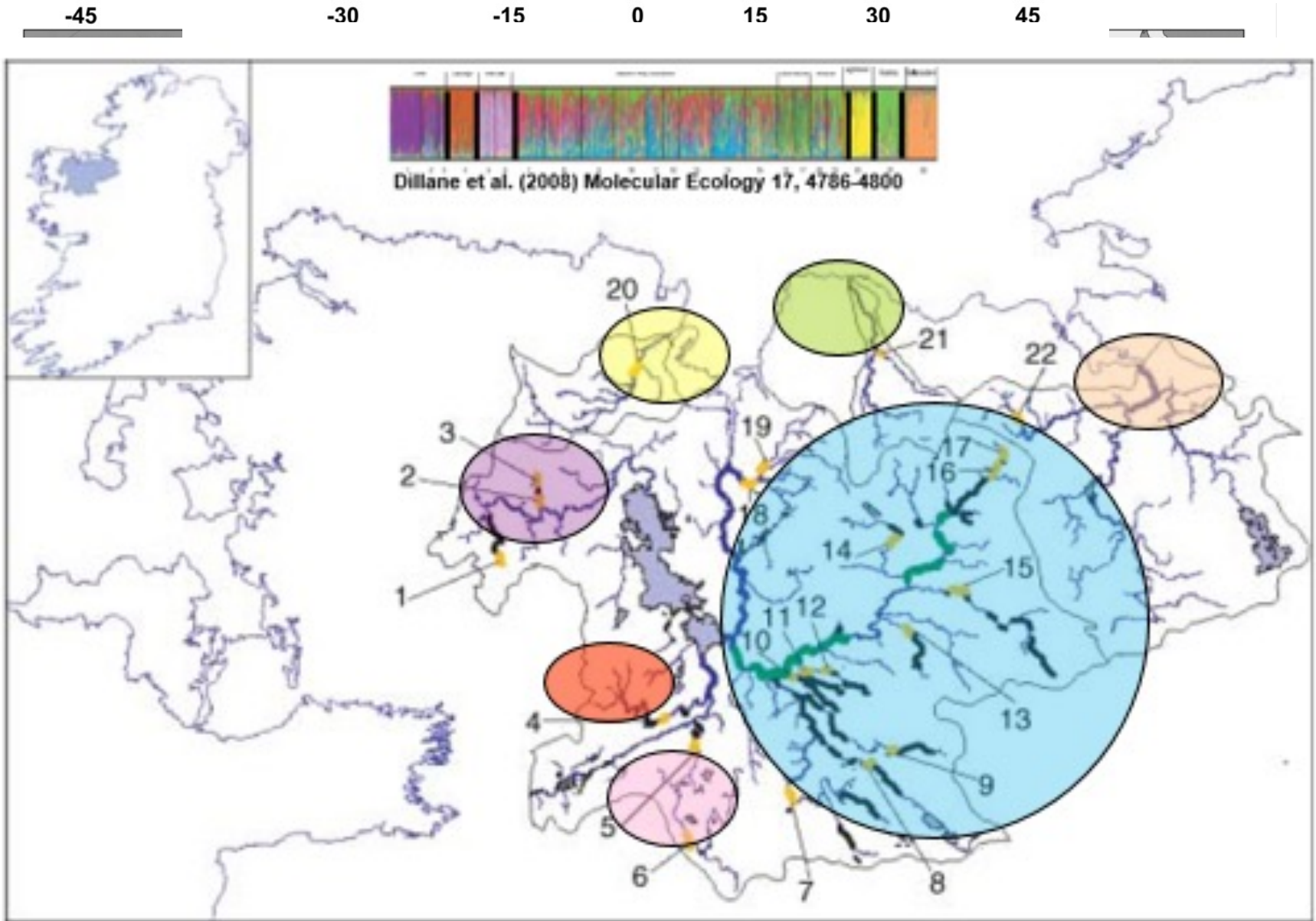
ICES Journal of Marine Science (2018), 75(2), 662–674. doi:10.1093/icesjms/txx184

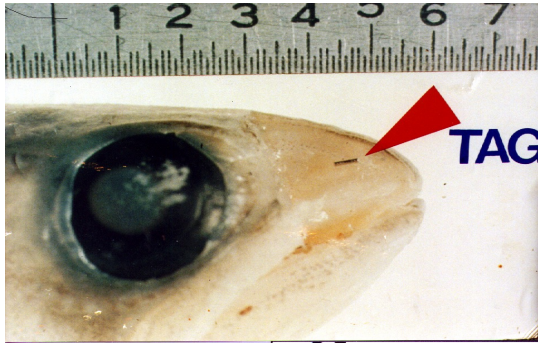
## Original Article

### A microsatellite baseline for genetic stock identification of European Atlantic salmon (*Salmo salar* L.)

John Gilbey<sup>1,\*</sup>, Jamie Coughlan<sup>2</sup>, Vidar Wennevik<sup>3</sup>, Paulo Prodöhl<sup>4</sup>, Jamie R. Stevens<sup>5</sup>, Carlos Garcia de Leaniz<sup>6</sup>, Dennis Ensing<sup>7</sup>, Eef Cauwelier<sup>1</sup>, Corrine Cherbonnel<sup>8</sup>, Sofia Consuegra<sup>6,9</sup>, Mark W. Coulson<sup>10,11</sup>, Tom F. Cross<sup>2</sup>, Walter Crozier<sup>7</sup>, Eileen Dillane<sup>2</sup>, Jonathan S. Ellis<sup>5,3</sup>, Eva García-Vázquez<sup>12</sup>, Andrew M. Griffiths<sup>5</sup>, Sigurdur Gudjonsson<sup>13</sup>, Kjetil Hindar<sup>14</sup>, Sten Karlsson<sup>14</sup>, David Knox<sup>1</sup>, Gonzalo Machado-Schiaffino<sup>12,5</sup>, Dorte Meldrup<sup>15</sup>, Einar Eg Nielsen<sup>15</sup>, Kristinn Ólafsson<sup>16</sup>, Craig R. Primmer<sup>17,8</sup>, Sergey Prusov<sup>18</sup>, Lee Stradmeyer<sup>1</sup>, Juha-Pekka Vähä<sup>17,\*,\*</sup>, Alexey Je. Veselov<sup>19</sup>, Lucy M. I. Webster<sup>10,11</sup>, Philip McGinnity<sup>2,1</sup>, and Eric Verspoor<sup>1,11,1</sup>

# Assignment Groups - Phylogenetic lineages





**Markers** (Analogous to physical markers)

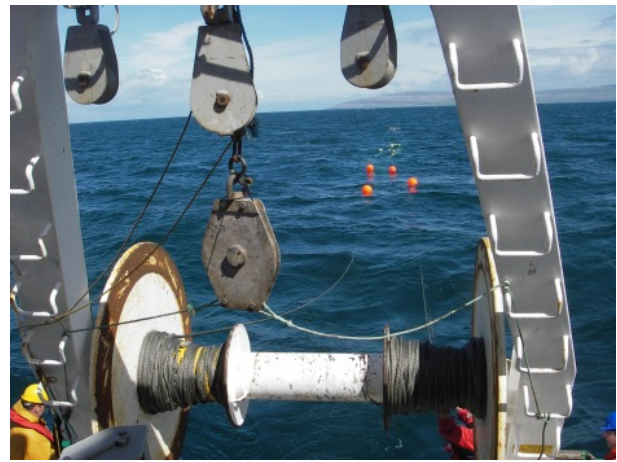
Statistically derived assignments rather than absolute – so quality of assignment will be a function of the quality of the baseline, which depends on coverage and molecular differentiation of the elements

Every fish collected (large numbers) delivers information (with geolocation)

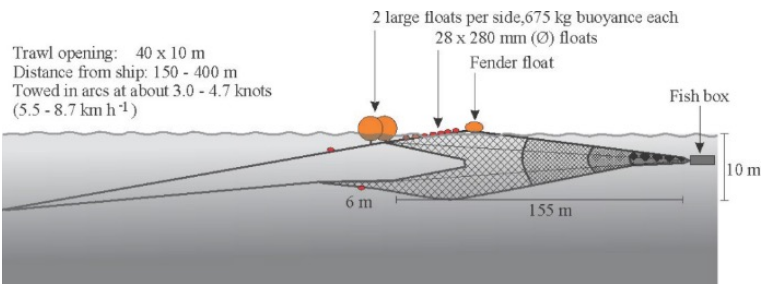
Other methods.....

## **Sampling post-smolts at sea**



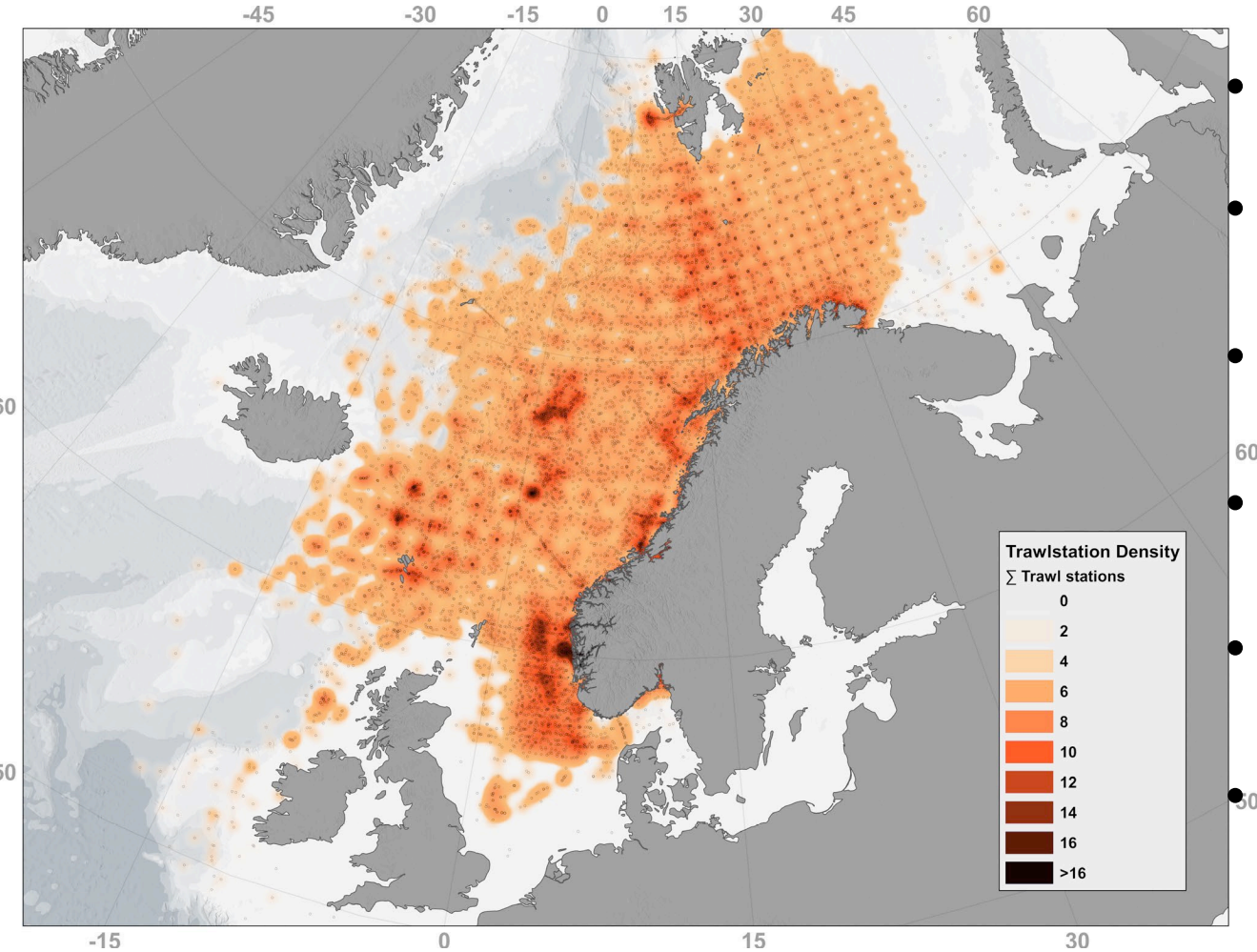


## Scientific pelagic cruises



# **Sampling coverage**

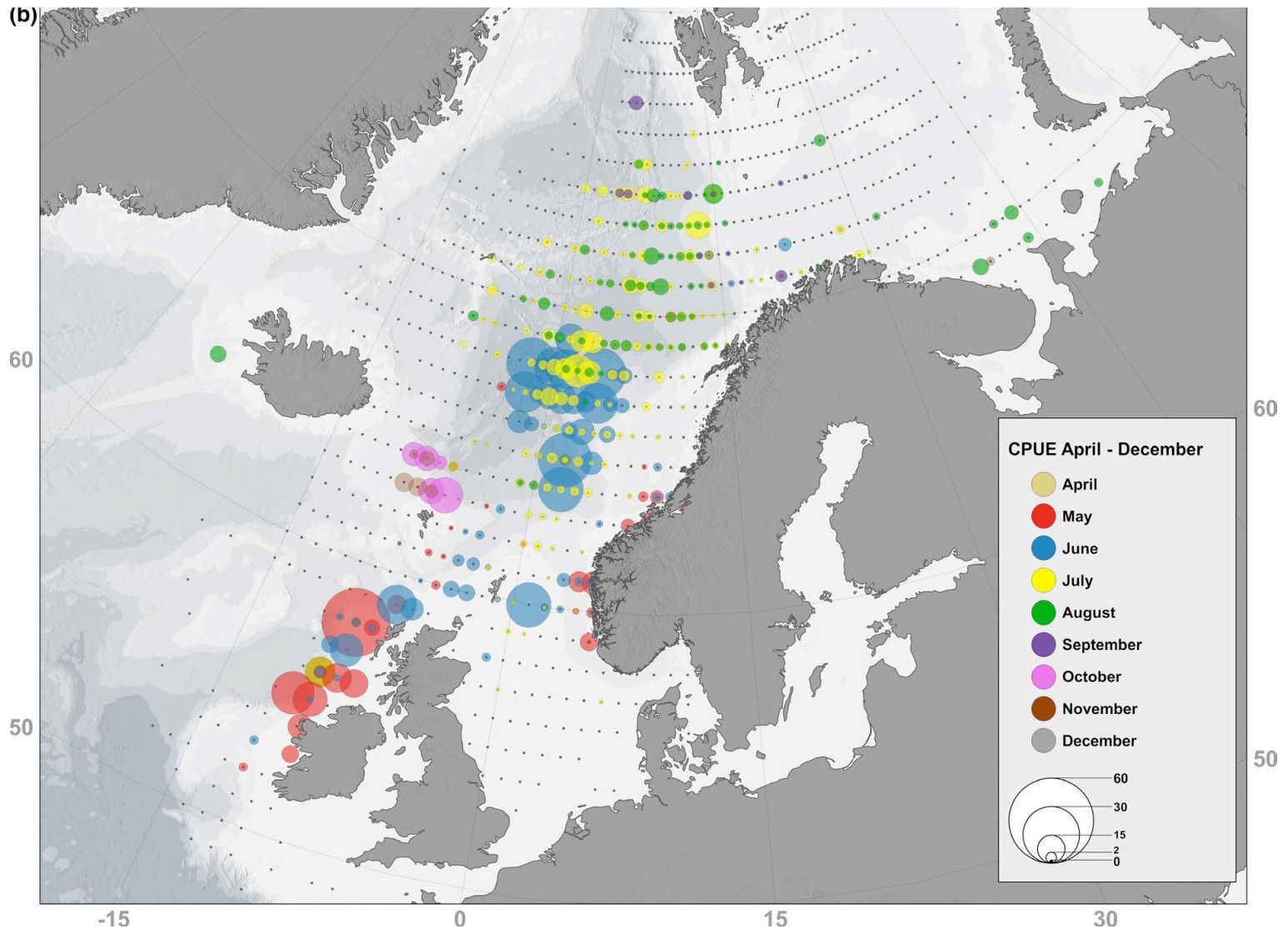
# Here's where the project took hold



- 4.75 million Km<sup>2</sup>
- 385 marine cruises
- 10,202 ind. trawls
- Sampled over 3 decades
- 9,269 post smolts
- 3,423 assigned to regional stock group

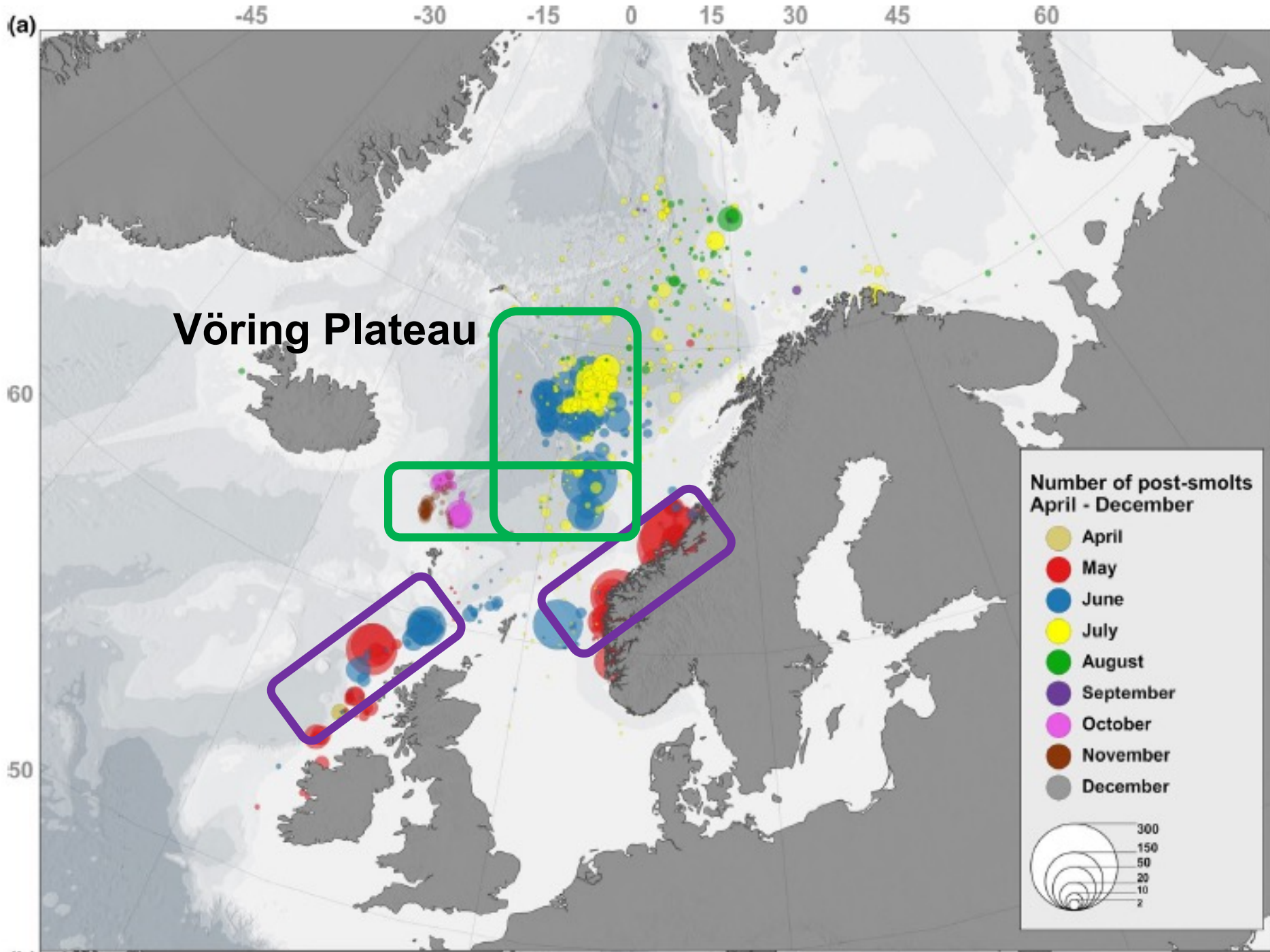
**Trawl-station density**

**Catch per unit effort (CPUE)**



**Key observation:** Where fish were ‘not found’ rather than where they were found

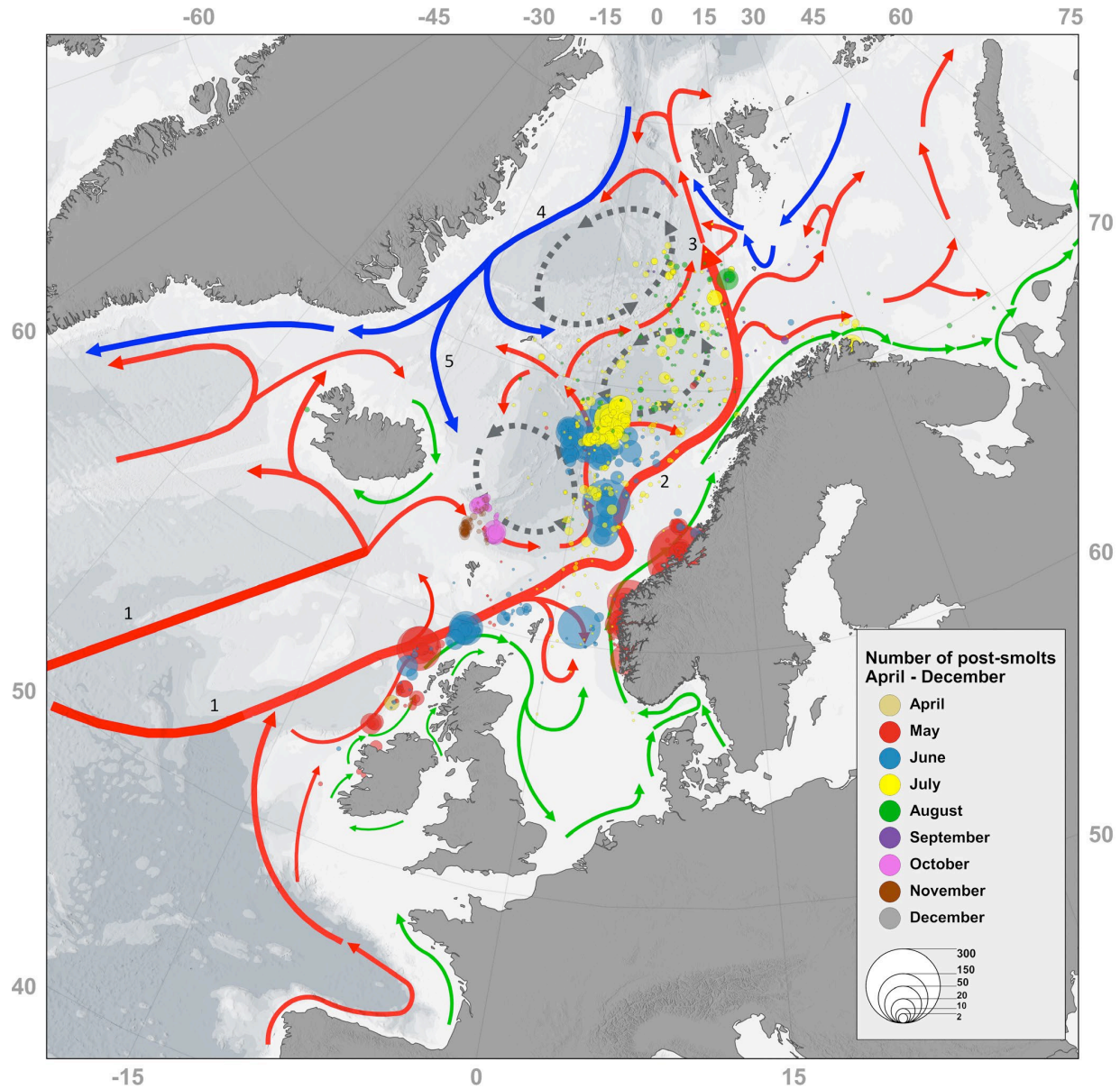
# **Migration routes & feeding aggregations**



# **Physiography (ocean habitat)**



# Seem to be able to get off the ocean current highway

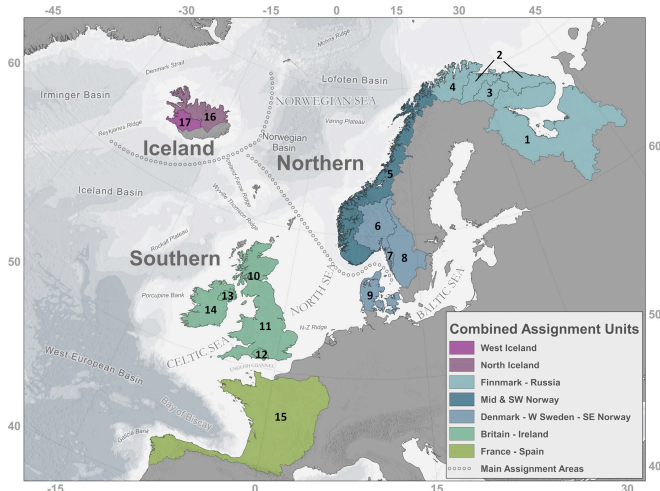


**Biological insights (stock specific ocean use)**

# What you expect to see v what you actually see (Observed v Expected)



## WORKING GROUP ON NORTH ATLANTIC SALMON (WGNAS)



Year	Northern NEAC					Northern NEAC (5%; 95%)	Southern NEAC					Southern NEAC (5%; 95%)	NEAC Area NEAC (5%; 95%)	
	Finland	Iceland (N&E)	Norway	Russia	Sweden		France	Iceland (S&W)	Ireland	UK(EW)	UK(NI)			UK(Scot)
Mean 10-year	32 124	18 282	262 295	123 661	6 521	446 925 (390 263; 514 320)	16 291	49 972	233 168	52 464	48 823	317 189	747 477 (596 070; 966 664)	1 198 029 (1 022 016; 1 435 065)

Table 3.3.4.3. Estimated pre-fishery abundance of maturing 1SW salmon (potential 1SW returns) by year for NEAC countries (50% quantile of the Monte Carlo distribution only) and region (50% (5%; 95%) quantiles of the Monte Carlo distribution).

Year	Northern NEAC					Northern NEAC (5%; 95%)	Southern NEAC					Southern NEAC (5%; 95%)	NEAC Area NEAC (5%; 95%)	
	Finland	Iceland (N&E)	Norway	Russia	Sweden		France	Iceland (S&W)	Ireland	UK(EW)	UK(NI)			UK(Scot)
Mean 10-year	30 012	8 926	411 962	115 225	15 435	584 080 (479 508; 711 449)	13 746	8 768	34 458	150 296	11 045	394 464	628 259 (450 087; 885 019)	1 216 996 (968 475; 1 539 364)

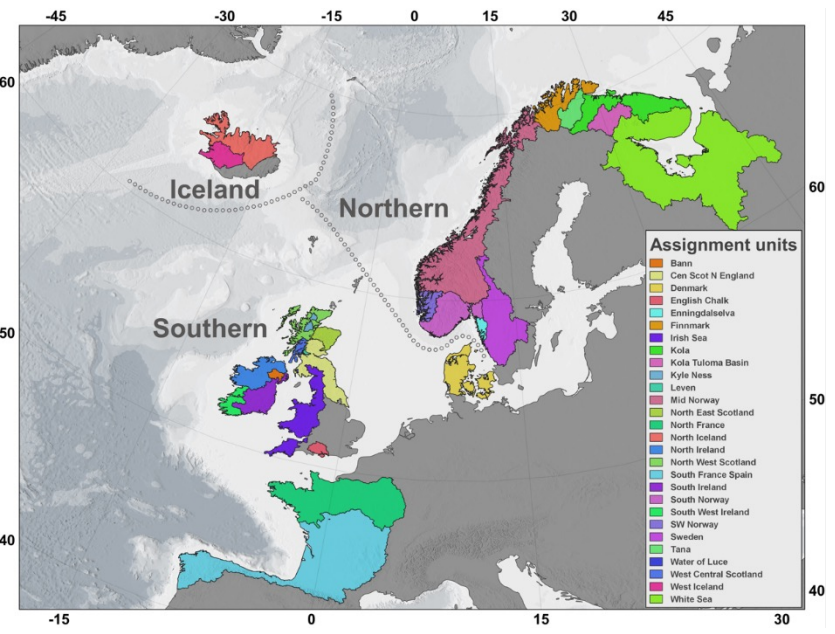
Table 3.3.4.4. Estimated pre-fishery abundance of non-maturing 1SW salmon (potential MSW returns) by year for NEAC countries (50% quantile of the Monte Carlo distribution only) and region (50% (5%; 95%) quantiles of the Monte Carlo distribution).

# Need higher resolution – can go back to individual river published catch (rod) data



Salmonid and Freshwater Fisheries Statistics for England and Wales, 2015  
 Including declared catches for salmon, sea trout, eels, smelt and lamprey by rods, nets and other instruments  
 February 2017

Environment Agency Salmon and freshwater fisheries statistics for England and Wales 2015 1



- > Labour market, earnings & education
- > Population & housing
- > Health & society
- > Environment & transport

Home / Agriculture, forestry, hunting and fishing / Fishing / Sea catches of salmon and sea trout

## Sea catches of salmon and sea trout

Updated: 25 November 2021

Next update: 25 November 2022

Change in sea catches  
 2019 - 2021  
**-27.2%**

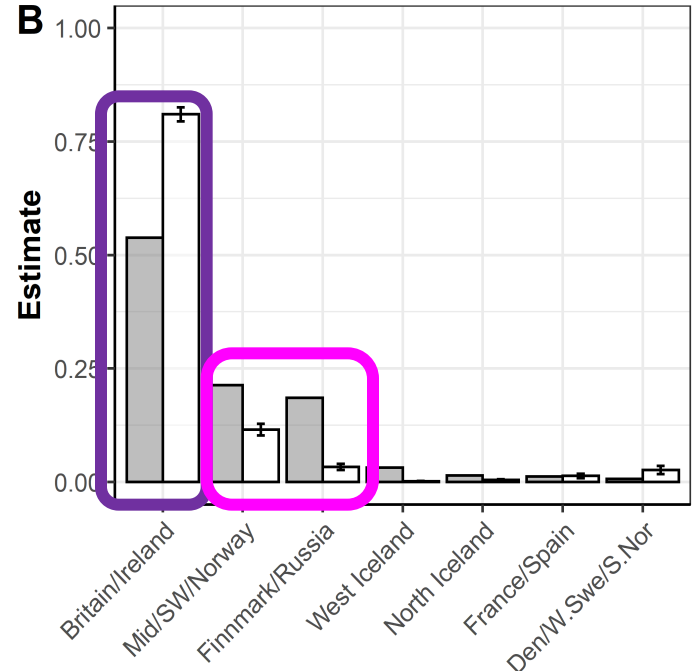
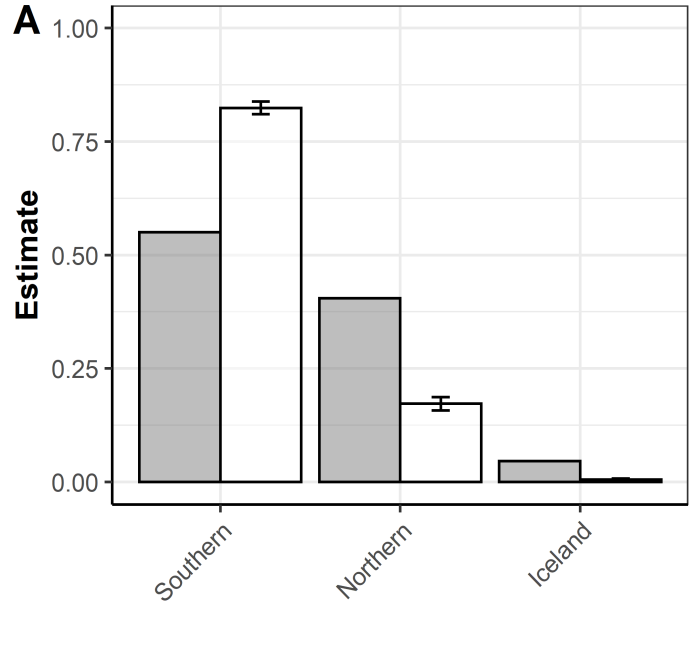
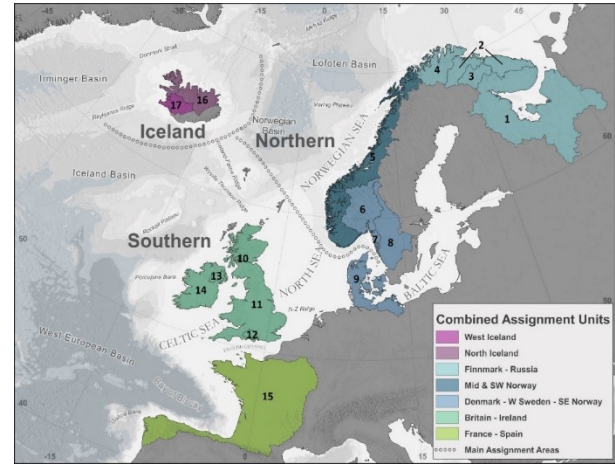
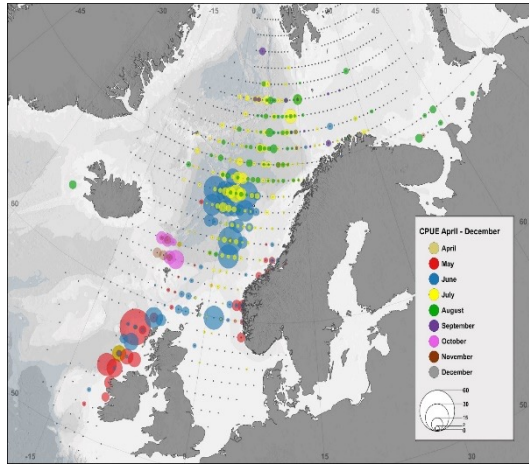
### Scottish Salmon and Sea Trout Fishery Statistics

- Official Statistics  
 The salmon and sea trout fishery statistics are published by Scottish Government in accordance with the Code of Practice for Official Statistics. Fishery statistics for a given season are published in the following April.



- Salmon Conservation Regulations  
 The Scottish Government has introduced a range of measures designed to improve the conservation status of wild salmon by managing their exploitation through fishing within Scotland's domestic waters. These regulations will have an effect on the catch and effort data reported by Scottish salmon fisheries.

# Some surprises at the stock level!!



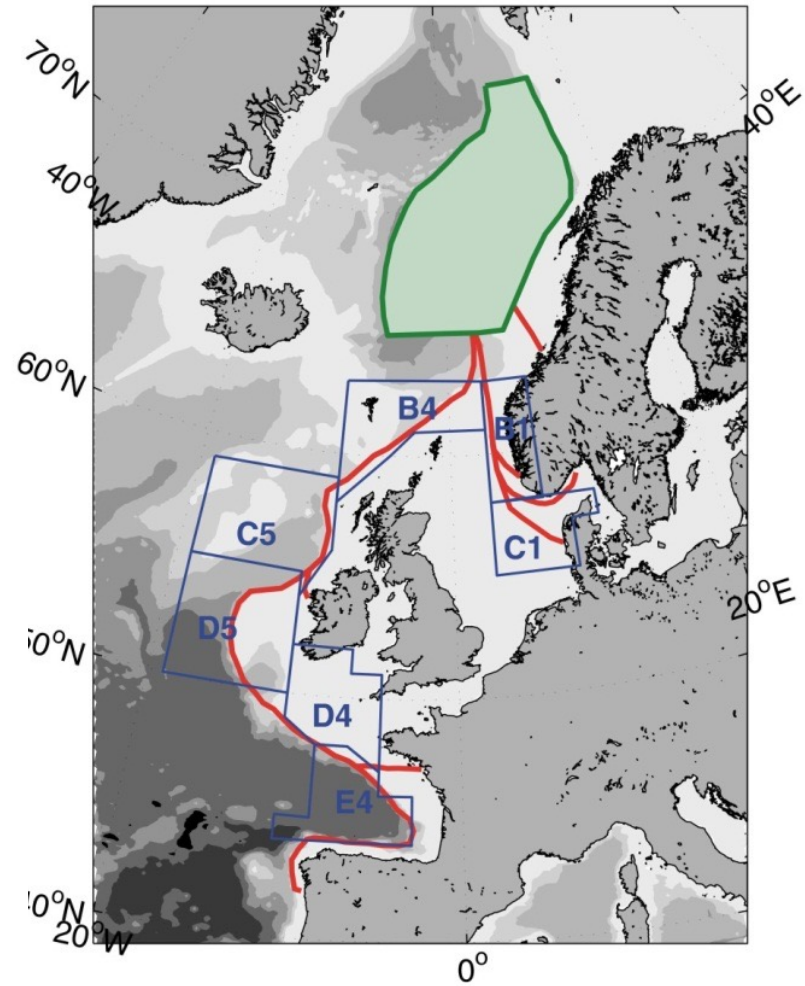
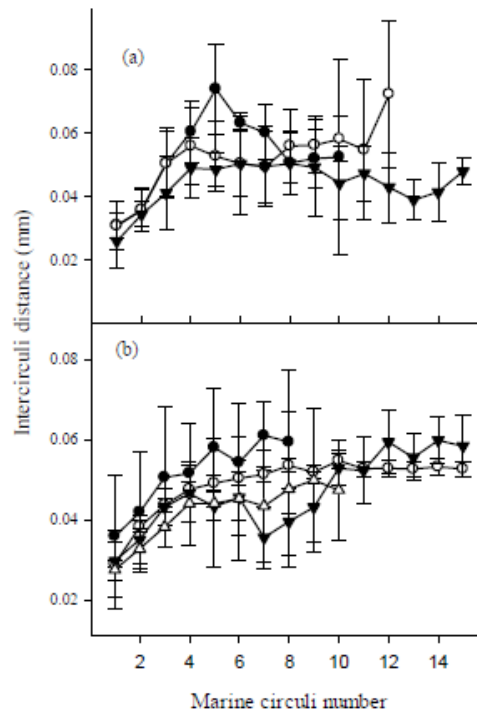
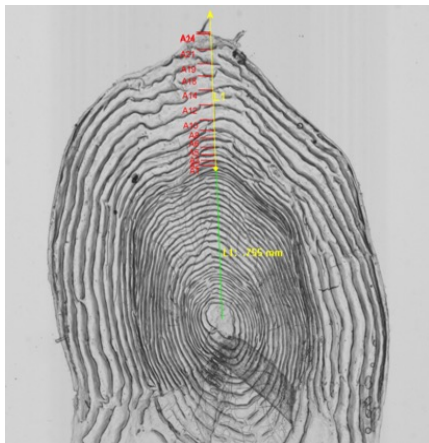
Estimate origin  ICES PFA  MSA

**Biological insights (survival proxies *in situ*)**

**When we look (work ongoing) at the stock groups we see *in situ* important stock specific differences in performance that might affect survival ultimately**

- Size at age
- Sex ratios
- Body condition
- Growth rates (acquired from scales)
- Feeding preferences
- Migration (swimming) speeds & distances covered

# Population specific growth rates



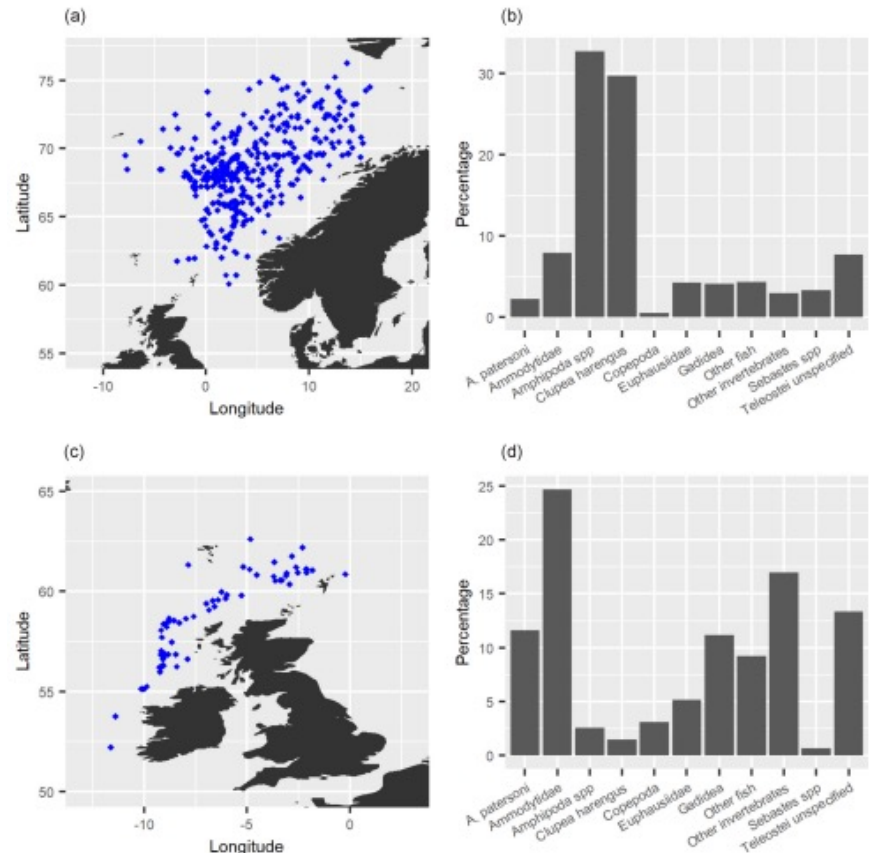


Original Article

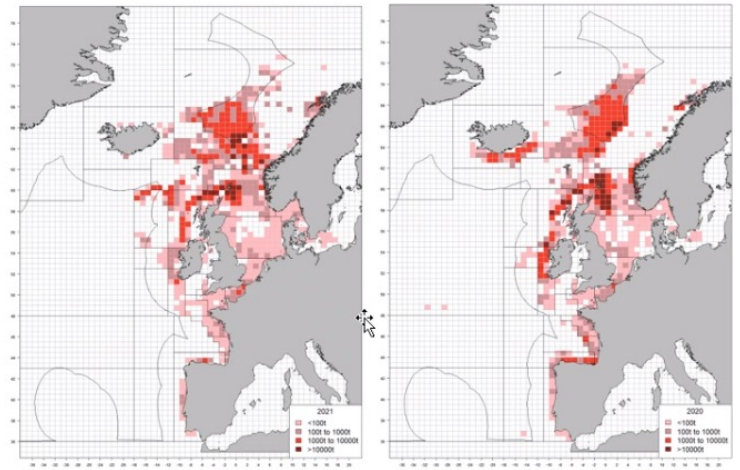
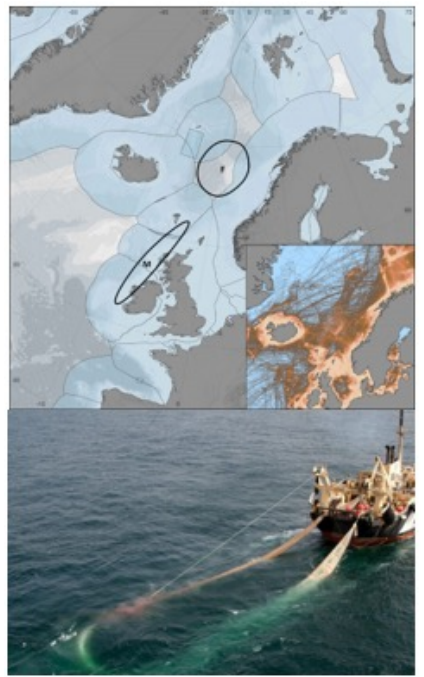
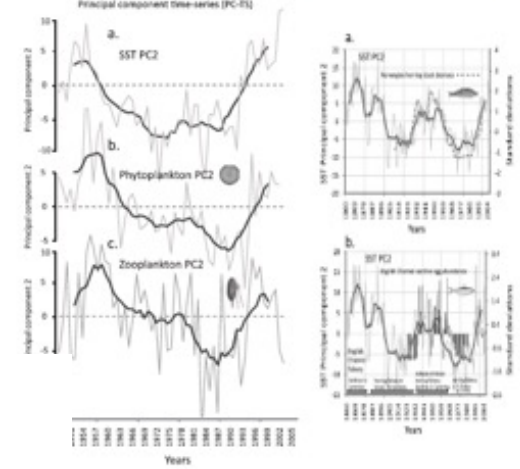
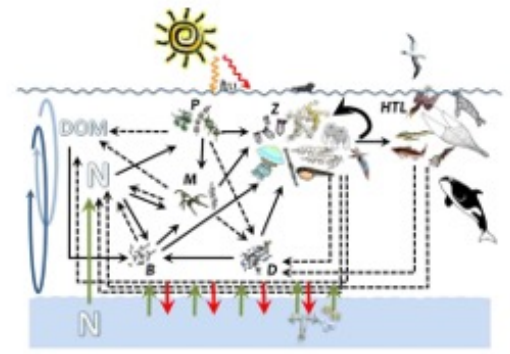
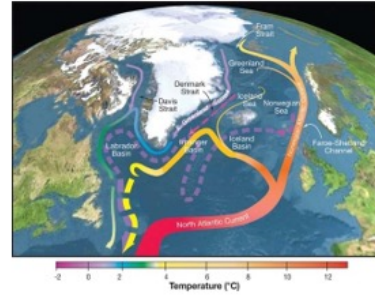
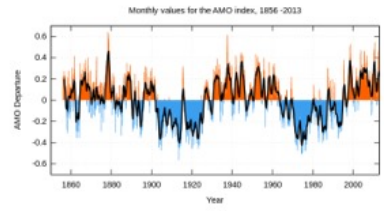
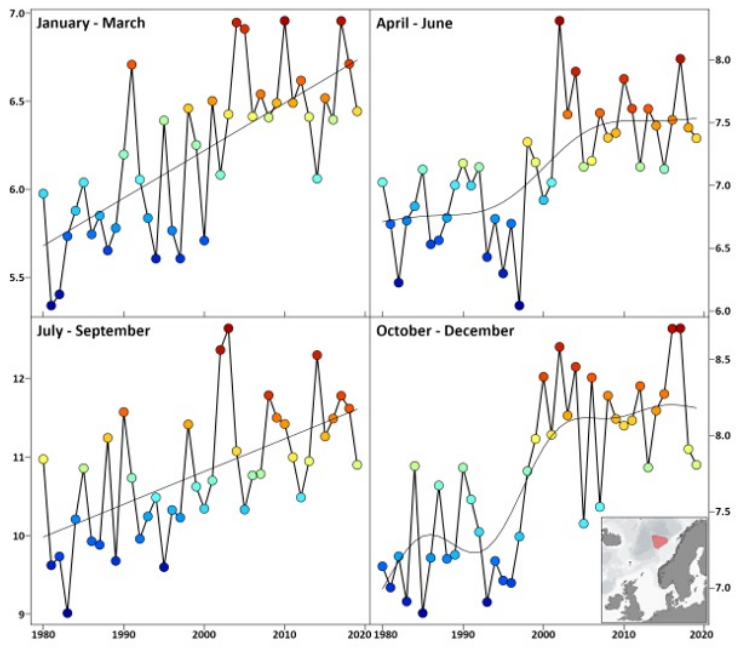
Poor feeding opportunities and reduced condition factor for salmon post-smolts in the Northeast Atlantic Ocean

Kjell Rong Utne<sup>1,\*</sup>, Beatriz Diaz Pauli<sup>2</sup>, Monika Haugland<sup>3</sup>, Jan Arge Jacobsen<sup>4</sup>, Niall Maoileidigh<sup>5</sup>, Webjørn Melle<sup>1</sup>, Cecilie Thorsen Broms<sup>1</sup>, Leif Nøttestad<sup>1</sup>, Marianne Holm<sup>1</sup>, Katie Thomas<sup>4</sup>, and Vidar Wennevik<sup>1</sup>

# Feeding preferences



**Biological (demographic) response (ecosystem variation)**



Drivers

# **The challenges of going beyond the Vöring Plateau**

# The challenges

- Locate migration paths and feeding grounds (European perspective) for Norwegian and Iceland post smolts – (possible solution – spatial and temporal strategic sampling required)
- Determine distribution of older fish over entire life history (possible solution – spatial and temporal strategic – **grid network of sampling stations** )
- Fish are growing rapidly once reach a certain size not easy to catch in pelagic trawls
- Migratory behaviour starts to differentiate on basis of trait development (e.g. maturation schedules/sex) – higher resolution baselines (possible solution)

# Excellent opportunistic studies

Fisheries Research 187 (2017) 110–119



Contents lists available at ScienceDirect

Fisheries Research

journal homepage: [www.elsevier.com/locate/fishres](http://www.elsevier.com/locate/fishres)



## Genetic stock identification of Atlantic salmon caught in the Faroese fishery

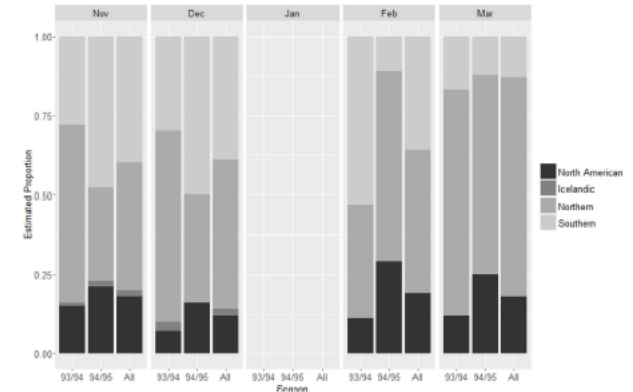
John Gilbey<sup>a,\*</sup>, Vidar Wennevik<sup>b</sup>, Ian R. Bradbury<sup>c</sup>, Peder Fiske<sup>d</sup>, Lars Petter Hansen<sup>d</sup>, Jan Arge Jacobsen<sup>e</sup>, Ted Potter<sup>f</sup>



Genetic stock identification reveals greater use of an oceanic feeding ground around the Faroe Islands by multi-sea winter Atlantic salmon.

Ronan James O'Sullivan<sup>1\*</sup>, Mikhail Ozerov<sup>2</sup>, Geir H. Bolstad<sup>3</sup>, John Gilbey<sup>4</sup>, Jan Arge Jacobsen<sup>5</sup>, Jaakko Erkinaro<sup>6</sup>, Audun H. Rikardsen<sup>6,7</sup>, Kjetil Hindar<sup>3</sup>, Tutku Aykanat<sup>1</sup>

J. Gilbey et al. / Fisheries Research 187 (2017) 110–119



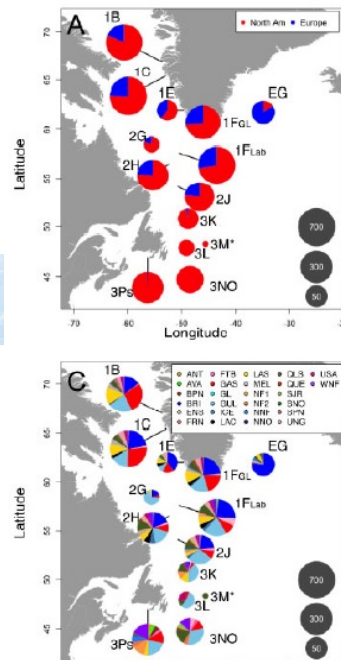
ICES Journal of Marine Science



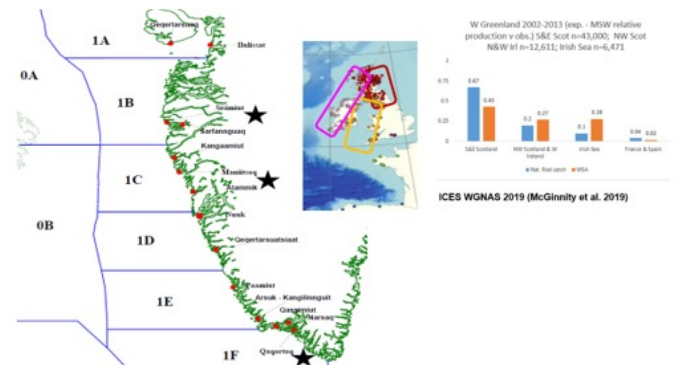
ICES Journal of Marine Science (2021), doi:10.1093/icesjms/fsaa152

## Range-wide genetic assignment confirms long-distance oceanic migration in Atlantic salmon over half a century

I. R. Bradbury<sup>1,2,\*</sup>, S. J. Lehnert<sup>1</sup>, A. Messmer<sup>1</sup>, S. J. Duffy<sup>1</sup>, E. Verspoor<sup>3</sup>, T. Kess<sup>1</sup>, J. Gilbey<sup>4</sup>, V. Wennevik<sup>5</sup>, M. Robertson<sup>1</sup>, G. Chaput<sup>6</sup>, T. Sheehan<sup>7</sup>, P. Bentzen<sup>3</sup>, J. B. Dempson<sup>1</sup>, and D. Reddin<sup>8</sup>



# New genetic markers / new baselines



# What makes a MSW salmon?

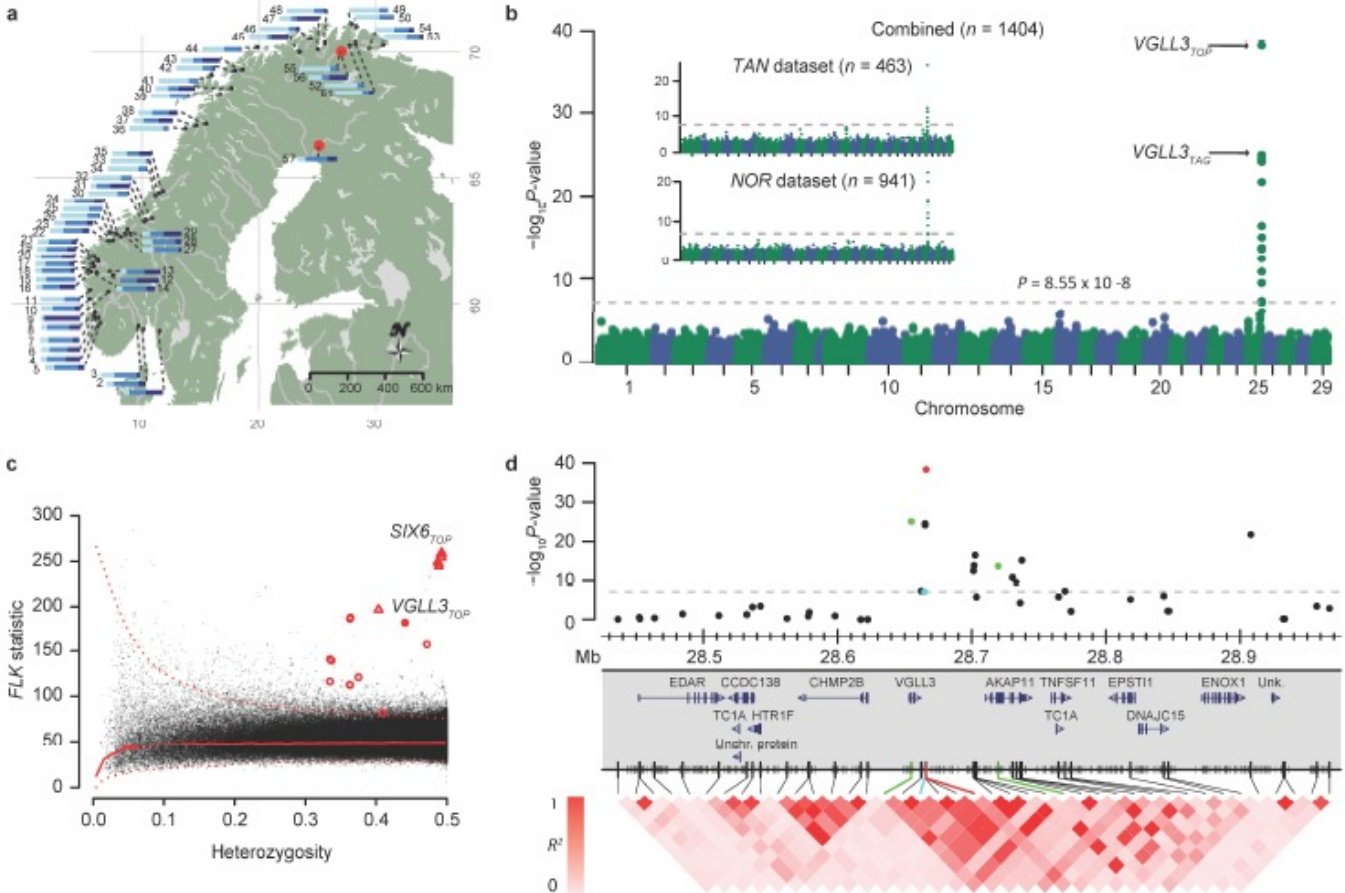
## LETTER

doi:10.1038/nature16062

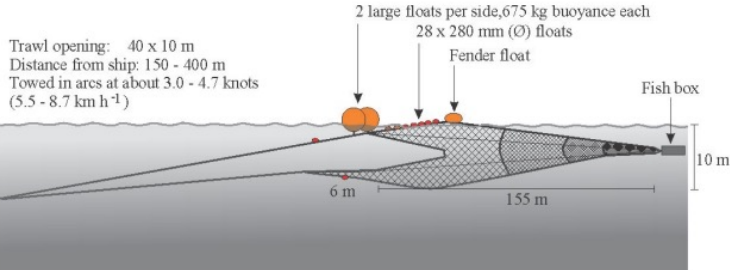
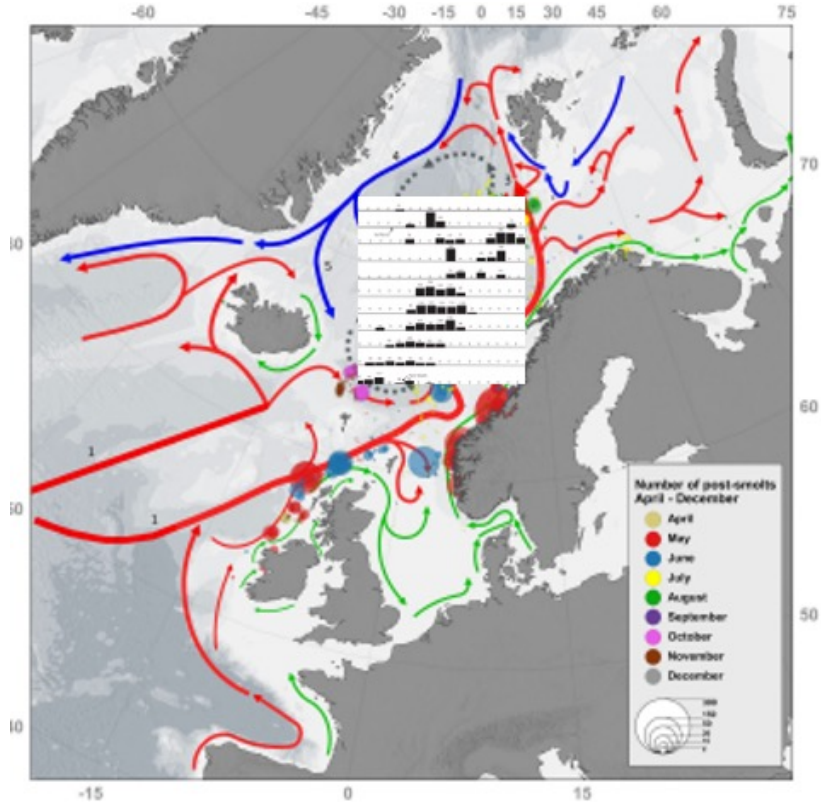
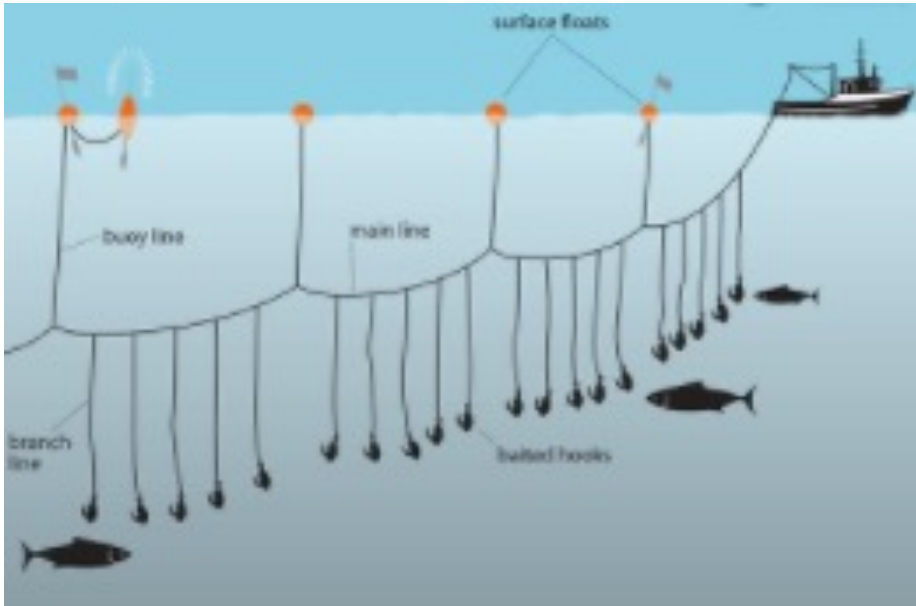
### Sex-dependent dominance at a single locus maintains variation in age at maturity in salmon

Nicola J. Barson<sup>1\*</sup>, Tutku Aykanat<sup>2\*</sup>, Kjetil Hindar<sup>3</sup>, Matthew Baranski<sup>4</sup>, Geir H. Bolstad<sup>3</sup>, Peder Fiske<sup>3</sup>, Céleste Jacq<sup>4</sup>, Arne J. Jensen<sup>3</sup>, Susan E. Johnston<sup>5</sup>, Sten Karlsson<sup>3</sup>, Matthew Kent<sup>1</sup>, Thomas Moen<sup>6</sup>, Eero Niemelä<sup>7</sup>, Torfinn Nome<sup>1</sup>, Tor F. Næsje<sup>3</sup>, Panu Orell<sup>7</sup>, Atso Romakkaniemi<sup>7</sup>, Harald Sægrov<sup>8</sup>, Kurt Urdal<sup>8</sup>, Jaakko Erkinaro<sup>7</sup>, Sigbjørn Lien<sup>1</sup> & Craig R. Primmer<sup>2</sup>

## VGLL3 locus



Jákupsstovu, S. H. I. (1988). Exploitation and migration of salmon in Faroese waters. In *Atlantic Salmon: Planning for the Future* (Mills, D. & Piggins, D., eds), pp. 458–482. London: Croom Helm.





# Concluding remark

Genetic stock Identification (marker) and genetics/genomics (evolutionary aspects) with strategic sampling has a powerful (central) contribution to make to our understanding of migration and distribution of Atlantic salmon in the sea

- *‘Advances in understanding are most likely to be realised by integrating insights from genetic-based distributional and telemetric studies and, given their respective limitations, using them to develop, parameterize and test migrational models. This integration will be the most powerful way to help define future conservation management challenges and priorities’ - from **Gilbey et al. (2021). Fish & Fisheries.***

